

RAW SEQUENCE LISTING

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: 10/506,763
Source: IFW16
Date Processed by STIC: 11/29/06

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IFW16

RAW SEQUENCE LISTING

DATE: 11/29/2006

PATENT APPLICATION: US/10/506,763

TIME: 09:19:32

Input Set : A:\Acceptys 0358749.ST25.txt

Output Set: N:\CRF4\11292006\J506763.raw

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3 <110> APPLICANT: ACCEPTYS, INC.
4     VOLLMERS , HEINZ P
5     MUELLER-HERMELINK , HANS K
7 <120> TITLE OF INVENTION: NEOPLASM SPECIFIC ANTIBODIES AND USES THEREOF
9 <130> FILE REFERENCE: 015065-0358749
11 <140> CURRENT APPLICATION NUMBER: 10/506,763
12 <141> CURRENT FILING DATE: 2005-05-12
14 <150> PRIOR APPLICATION NUMBER: DE 102 10 427.1
15 <151> PRIOR FILING DATE: 2002-03-09
17 <150> PRIOR APPLICATION NUMBER: PCT/IB2003/01335
18 <151> PRIOR FILING DATE: 2003-03-10
20 <160> NUMBER OF SEQ ID NOS: 4
22 <170> SOFTWARE: PatentIn version 3.3
24 <210> SEQ ID NO: 1
25 <211> LENGTH: 102
26 <212> TYPE: PRT
27 <213> ORGANISM: Homo sapiens
30 <220> FEATURE:
31 <221> NAME/KEY: misc_feature
32 <223> OTHER INFORMATION: Heavy chain CM-1 variable region
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41           20           25           30
44 Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala Asp Ser
45           35           40           45
48 Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr Leu
49           50           55           60
52 Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr Tyr
53 65           70           75           80
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65 <211> LENGTH: 306
66 <212> TYPE: DNA
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70 <220> FEATURE:
71 <221> NAME/KEY: misc_feature
72 <223> OTHER INFORMATION: Heavy chain CM-1 variable region
74 <400> SEQUENCE: 2

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75 aggtccctga gactctcctg tgcagcctct ggattcacct tcagtagcta tggcatgcac      60
77 tgggtccgcc aggtccagg caaggggctg gagtgggtgg cagttatatc atatgatgga      120
79 agtaataaat actatgcaga ctccgtgaag ggccgattca ccatctccag agacaattcc      180
81 aagaacacgc tgtatctgca aatgaacagc ctgagagctg aggacacggc tgtgtattac      240
83 tgtgcgaaag accggtcttc gggctactac ggtatggacg tctggggcca aggcaccctg      300
85 gtcacc                                           306
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89 <211> LENGTH: 109
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104 Thr Ala Arg Ile Thr Cys Ser Gly Asp Ala Leu Pro Lys Gln Tyr Ala
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108 Tyr Trp Tyr Gln Gln Lys Pro Gly Gln Ala Pro Val Leu Val Ile Tyr
109      35                    40                    45
112 Lys Asp Ser Glu Arg Pro Ser Gly Ile Pro Glu Arg Phe Ser Gly Ser
113      50                    55                    60
116 Ser Ser Gly Thr Thr Val Thr Leu Thr Ile Ser Gly Val Gln Ala Glu
117 65                    70                    75                    80
120 Asp Glu Ala Asp Tyr Tyr Cys Gln Ser Ala Asp Ser Ser Gly Thr Tyr
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134 <220> FEATURE:
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143 caggccccctg tgctgggtgat atataaagac agtgagaggc cctcagggat ccctgagcga      180
145 ttctctggct ccagctcagg gacaacagtc acgttgacca tcagtggagt ccaggcagaa      240
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149 ggagggacca agctgaccgt cctaggt                                           327

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VERIFICATION SUMMARY

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